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SEQUENCE LISTING

<110> Japan Science and Technology Corporation

<120> Diagnosis method of endometriosis

<130> 03-F-104PCT

<150> JP2003-196459

<151> 2003-07-14

<160> 10

<170> PatentIn version 3.1

<210> 1

<211> 830

<212> DNA

<213> Homo sapiens

<220>

<221> CDS

<222> (95)..(613)

<223>

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 Met Ile Ile Tyr Arg Asp Leu
 1 5

atc agc cac gat gag atg ttc tcc gac atc tac aag atc cgg gag atc 163
 Ile Ser His Asp Glu Met Phe Ser Asp Ile Tyr Lys Ile Arg Glu Ile
 10 15 20

gcg gac ggg ttg tgc ctg gag gtg gag ggg aag atg gtc agt agg aca 211
 Ala Asp Gly Leu Cys Leu Glu Val Glu Gly Lys Met Val Ser Arg Thr
 25 30 35

gaa ggt aac att gat gac tcg ctc atl ggt gga aat gcc tcc gct gaa 259
 Glu Gly Asn Ile Asp Asp Ser Leu Ile Gly Gly Asn Ala Ser Ala Glu
 40 45 50 55

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ggc ccc gag ggc gaa ggt acc gac aca gta atc act ggt gtc gat Gly Pro Glu Gly Glu Gly Thr Glu Ser Thr Val Ile Thr Gly Val Asp 60 65 70	307
att gtc atg aac cat cac ctg cag gaa aca agt ttc aca aaa gaa gcc Ile Val Met Asn His His Leu Gln Glu Thr Ser Phe Thr Lys Glu Ala 75 80 85	355
tac aag aag tac atc aaa gat tac atg aaa tca atc aaa ggg aaa ctt Tyr Lys Lys Tyr Ile Lys Asp Tyr Met Lys Ser Ile Lys Gly Lys Leu 90 95 100	403
gaa gaa cag aga cca gaa aga gta aaa cct ttt atg aca ggg gct gca Glu Glu Gln Arg Pro Glu Arg Val Lys Pro Phe Met Thr Gly Ala Ala 105 110 115	451
gaa caa atc aag cac atc ctt gct aat ttc aaa aac tac cag ttc ttt Glu Gln Ile Lys His Ile Leu Ala Asn Phe Lys Asn Tyr Gln Phe Phe 120 125 130 135	499
att ggt gaa aac atg aat cca gat ggc atg gtt gct cta ttg gac tac Ile Gly Glu Asn Met Asn Pro Asp Gly Met Val Ala Leu Leu Asp Tyr 140 145 150	547
cgt gag gai ggt gtg acc cca tat atg att ttc ttt aag gat ggt tta Arg Glu Asp Gly Val Thr Pro Tyr Met Ile Phe Phe Lys Asp Gly Leu 155 160 165	595
gaa atg gaa aaa tgt taa caaatgtggc aattatttt gatctatcac Glu Met Glu Lys Cys 170	643
cgtcatcat aactggcttc tgcttgtcat ccacacaaca ccaggactta agacaaatgg 703	
gacttgtatgc accttgatgt ctccatttat ttgtactgtg attttttgg agggaggca 763	
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<210> 2
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<212> PRT

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<213> Homo sapiens

<400> 2

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Gly Lys Met Val Ser Arg Thr Glu Gly Asn Ile Asp Asp Ser Leu Ile			
35	40	45	
Gly Gly Asn Ala Ser Ala Glu Gly Pro Glu Gly Glu Gly Thr Glu Ser			
50	55	60	
Thr Val Ile Thr Gly Val Asp Ile Val Met Asn His His Leu Gln Glu			
65	70	75	80
Thr Ser Phe Thr Lys Glu Ala Tyr Lys Lys Tyr Ile Lys Asp Tyr Met			
85	90	95	
Lys Ser Ile Lys Gly Lys Leu Glu Glu Arg Pro Glu Arg Val Lys			
100	105	110	
Pro Phe Met Thr Gly Ala Ala Glu Gln Ile Lys His Ile Leu Ala Asn			
115	120	125	
Phe Lys Asn Tyr Gln Phe Phe Ile Gly Glu Asn Met Asn Pro Asp Gly			
130	135	140	
Met Val Ala Leu Leu Asp Tyr Arg Glu Asp Gly Val Thr Pro Tyr Met			
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Ile Phe Phe Lys Asp Gly Leu Glu Met Glu Lys Cys			
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<212> DNA

<213> Artificial

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<223> Description of Artificial sequence: Synthetic oligonucleotide

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<210> 4

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<213> Artificial

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<210> 5

<211> 23

<212> DNA

<213> Artificial

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<212> DNA

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<223> Description of Artificial sequence: Synthetic oligonucleotide

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<210> 8
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<212> DNA
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<210> 9
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<210> 10
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<212> DNA
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<220>
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<400> 10
ggccgaattc agatccaaaa taattgcc

28

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